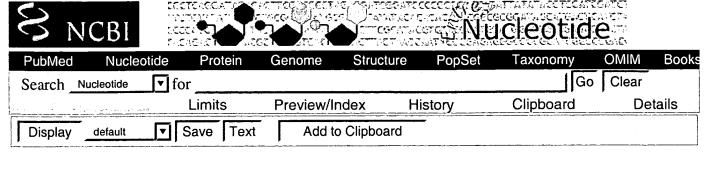


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Revised: October 24, 2001.



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Related Sequences, OMIM, Protein, PubMed, Taxonomy, UniSTS, LinkOut

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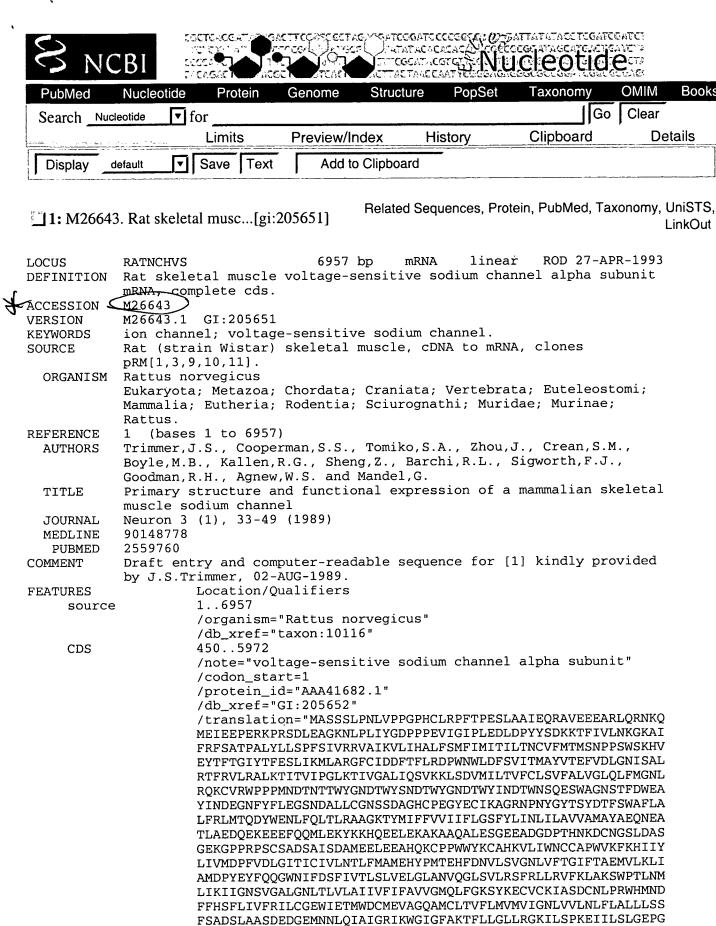
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Revised: October 24, 2001.

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1: P15389. SODIUM CHANNEL PR...[gi:116452]

BLink, Related Sequences, PubMed, Taxonomy, LinkOut

ROD 15-DEC-1998 2019 aa linear LOCUS CIN5_RAT SODIUM CHANNEL PROTEIN, CARDIAC MUSCLE ALPHA-SUBUNIT. DEFINITION ACCESSION (P15389

g116452 PID

VERSION P15389 GI:116452

swissprot: locus CIN5_RAT, accession P15389; **DBSOURCE**

class: standard. created: Apr 1, 1990.

sequence updated: Apr 1, 1990. annotation updated: Dec 15, 1998.

xrefs: gi: 206857, gi: 206858, gi: 112312

xrefs (non-sequence databases): PFAM PF00520, PFAM PF00612 KEYWORDS

Ionic channel; Transmembrane; Ion transport; Voltage-gated channel; Glycoprotein; Duplication; Multigene family; Phosphorylation.

SOURCE Norway rat.

ORGANISM Rattus norvegicus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

Rattus.

(residues 1 to 2019) REFERENCE

Rogart, R.B., Cribbs, L.L., Muglia, L.K., Kephart, D.D. and Kaiser, M.W. AUTHORS TITLE

Molecular cloning of a putative tetrodotoxin-resistant rat heart

Na+ channel isoform

Proc. Natl. Acad. Sci. U.S.A. 86 (20), 8170-8174 (1989) JOURNAL

MEDLINE 90046760 PUBMED 2554302

REMARK SEQUENCE FROM N.A.

TISSUE=HEART

COMMENT

This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. The original entry is available from http://www.expasy.ch/sprot and http://www.ebi.ac.uk/sprot

[FUNCTION] THIS PROTEIN MEDIATES THE VOLTAGE-DEPENDENT SODIUM ION PERMEABILITY OF EXCITABLE MEMBRANES. ASSUMING OPENED OR CLOSED CONFORMATIONS IN RESPONSE TO THE VOLTAGE DIFFERENCE ACROSS THE MEMBRANE, THE PROTEIN FORMS A SODIUM-SELECTIVE CHANNEL THROUGH WHICH NA+ IONS MAY PASS IN ACCORDANCE WITH THEIR ELECTROCHEMICAL GRADIENT. IT IS A TETRODOTOXIN-RESISTANT NA+ CHANNEL ISOFORM. [SUBCELLULAR LOCATION] INTEGRAL MEMBRANE PROTEIN.

[DOMAIN] THE SEQUENCE CONTAINS 4 INTERNAL REPEATS, EACH WITH 5 HYDROPHOBIC SEGMENTS (S1,S2,S3,S5,S6) AND ONE POSITIVELY CHARGED SEGMENT (S4). SEGMENTS S4 ARE PROBABLY THE VOLTAGE-SENSORS AND ARE CHARACTERIZED BY A SERIES OF POSITIVELY CHARGED AMINO ACIDS AT EVERY THIRD POSITION.

[MISCELLANEOUS] NA+ CHANNELS IN MAMMALIAN CARDIAC MEMBRANE HAVE FUNCTIONAL PROPERTIES QUITE DISTINCT FROM NA+ CHANNELS IN NERVE AND SKELETAL MUSCLE.

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[SIMILARITY] TO OTHER SODIUM CHANNEL PROTEINS.
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5/11/02 2:59 PM



PubMed	Nucleotide	Protein	Genome	Structure	PopSet	Taxonomy	OMIM	Books
Search Pro	otein ▼ fc	or				Go	Clear	
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1: Q99250. Sodium channel pr...[gi:6648080]

BLink, OMIM, Related Sequences, PubMed, Taxonomy, LinkOut

LOCUS 2005 aa linear PRI 01-MAR-2002 CIN2_HUMAN Sodium channel protein, brain II alpha subunit. DEFINITION ACCESSION Q99250 g6648080 PID Q99250 GI:6648080 VERSION swissprot: locus CIN2_HUMAN, accession Q99250; DBSOURCE class: standard. extra accessions:Q14472,created: Jun 1, 1994. sequence updated: May 30, 2000. annotation updated: Mar 1, 2002. xrefs: gi: 456678, gi: 457879, gi: 3075512, gi: 3075513, gi: 36419, gi: 36420, gi: 338282, gi: 179560, gi: 12750754, gi: 418893 xrefs (non-sequence databases): MIM 182390, InterPro IPR001682, InterPro IPR002111, InterPro IPR000048, InterPro IPR000636, InterPro IPR001696, Pfam PF00520, Pfam PF00612, PRINTS PR00170, SMART SM00015, PROSITE PS50096 **KEYWORDS** Ionic channel; Transmembrane; Ion transport; Voltage-gated channel; Glycoprotein; Repeat; Multigene family. SOURCE human. ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. REFERENCE (residues 1 to 2005) **AUTHORS** Ahmed, C.M., Ware, D.H., Lee, S.C., Patten, C.D., Ferrer-Montiel, A.V., Schinder, A.F., McPherson, J.D., Wagner-Mcpherson, C.B., Wasmuth, J.J., Evans, G.A. and Montal, M. TITLE Primary structure, chromosomal localization, and functional expression of a voltage-gated sodium channel from human brain JOURNAL Proc. Natl. Acad. Sci. U.S.A. 89 (17), 8220-8224 (1992) MEDLINE 92390418 REMARK SEQUENCE FROM N.A. TISSUE=Brain REFERENCE (residues 1 to 2005) **AUTHORS** Lu, C.-M., Eichelberger, J.S., Beckman, M.L., Schade, S.D. and Brown, G.B. TITLE Direct Submission JOURNAL Submitted (~APR-1998) SEQUENCE OF 1-89 FROM N.A. REMARK REFERENCE (residues 1 to 2005) **AUTHORS** Lu, C.M., Han, J., Rado, T.A. and Brown, G.B. TITLE Differential expression of two sodium channel subtypes in human brain **JOURNAL** FEBS Lett. 303 (1), 53-58 (1992) MEDLINE 92275082 REMARK SEQUENCE OF 1702-2005 FROM N.A. TISSUE=Brain REFERENCE (residues 1 to 2005) **AUTHORS** Han, J.A., Lu, C.M., Brown, G.B. and Rado, T.A.

Direct amplification of a single dissected chromosomal segment by polymerase chain reaction: a human brain sodium channel gene is on

TITLE

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            Proc. Natl. Acad. Sci. U.S.A. 88 (2), 335-339 (1991)
  JOURNAL
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COMMENT
            On Dec 30, 1999 this sequence version replaced gi:544037.
            This SWISS-PROT entry is copyright. It is produced through a
            collaboration between the Swiss Institute of Bioinformatics and
            the EMBL outstation - the European Bioinformatics Institute.
            The original entry is available from http://www.expasy.ch/sprot
            and http://www.ebi.ac.uk/sprot
            [FUNCTION] THIS PROTEIN MEDIATES THE VOLTAGE-DEPENDENT SODIUM ION
            PERMEABILITY OF EXCITABLE MEMBRANES. ASSUMING OPENED OR CLOSED
            CONFORMATIONS IN RESPONSE TO THE VOLTAGE DIFFERENCE ACROSS THE
            MEMBRANE, THE PROTEIN FORMS A SODIUM-SELECTIVE CHANNEL THROUGH
            WHICH NA++ IONS MAY PASS IN ACCORDANCE WITH THEIR ELECTROCHEMICAL
            GRADIENT.
            [SUBUNIT] THE SODIUM CHANNEL CONSISTS OF A LARGE POLYPEPTIDE AND
            2-3 SMALLER ONES. THIS SEQUENCE REPRESENTS A LARGE POLYPEPTIDE.
            [SUBCELLULAR LOCATION] Integral membrane protein.
            [DOMAIN] THE SEQUENCE CONTAINS 4 INTERNAL REPEATS, EACH WITH 5
            HYDROPHOBIC SEGMENTS (S1,S2,S3,S5,S6) AND ONE POSITIVELY CHARGED
            SEGMENT (S4). SEGMENTS S4 ARE PROBABLY THE VOLTAGE-SENSORS AND ARE
            CHARACTERIZED BY A SERIES OF POSITIVELY CHARGED AMINO ACIDS AT
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6 of 6 5/11/02 2:59 PM

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Search Pro	otein 🔽 fo	or				Go	Clear	
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1: P04775. Sodium channel pr...[gi:116448]

BLink, Related Sequences, PubMed, Taxonomy, LinkOut

CIN2_RAT 2005 aa linear ROD 16-OCT-2001 LOCUS DEFINITION Sodium channel protein, brain II alpha subunit. P04775

ACCESSION PTD

g116448

VERSION

(P04775) GI:116448

DBSOURCE

swissprot: locus CIN2_RAT, accession P04775;

class: standard.

created: Aug 13, 1987.

sequence updated: Aug 13, 1987. annotation updated: Oct 16, 2001. xrefs: gi: 57214, gi: 57215, gi: 92753

xrefs (non-sequence databases): InterPro IPR002111, InterPro IPR000636, InterPro IPR001682, InterPro IPR000048, InterPro IPR001696, Pfam PF00520, Pfam PF00612, PRINTS PR00170, SMART

SM00015, PROSITE PS50096

KEYWORDS

Ionic channel; Transmembrane; Ion transport; Voltage-gated channel;

Glycoprotein; Repeat; Multigene family.

SOURCE

Norway rat.

ORGANISM Rattus norvegicus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

Rattus.

REFERENCE

(residues 1 to 2005)

AUTHORS Noda, M., Ikeda, T., Kayano, T., Suzuki, H., Takeshima, H., Kurasaki, M.,

Takahashi, H. and Numa, S.

TITLE JOURNAL Existence of distinct sodium channel messenger RNAs in rat brain

Nature 320 (6058), 188-192 (1986)

MEDLINE 86146901

REMARK

SEQUENCE FROM N.A.

COMMENT

This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. The original entry is available from http://www.expasy.ch/sprot and http://www.ebi.ac.uk/sprot

[FUNCTION] THIS PROTEIN MEDIATES THE VOLTAGE-DEPENDENT SODIUM ION PERMEABILITY OF EXCITABLE MEMBRANES. ASSUMING OPENED OR CLOSED CONFORMATIONS IN RESPONSE TO THE VOLTAGE DIFFERENCE ACROSS THE MEMBRANE, THE PROTEIN FORMS A SODIUM-SELECTIVE CHANNEL THROUGH WHICH NA++ IONS MAY PASS IN ACCORDANCE WITH THEIR ELECTROCHEMICAL GRADIENT.

[SUBUNIT] THE SODIUM CHANNEL CONSISTS OF A LARGE POLYPEPTIDE AND 2-3 SMALLER ONES. THIS SEQUENCE REPRESENTS A LARGE POLYPEPTIDE. [SUBCELLULAR LOCATION] INTEGRAL MEMBRANE PROTEIN.

[DOMAIN] THE SEQUENCE CONTAINS 4 INTERNAL REPEATS, EACH WITH 5 HYDROPHOBIC SEGMENTS (S1,S2,S3,S5,S6) AND ONE POSITIVELY CHARGED SEGMENT (S4). SEGMENTS S4 ARE PROBABLY THE VOLTAGE-SENSORS AND ARE CHARACTERIZED BY A SERIES OF POSITIVELY CHARGED AMINO ACIDS AT EVERY THIRD POSITION.

[SIMILARITY] TO OTHER SODIUM CHANNEL PROTEINS.

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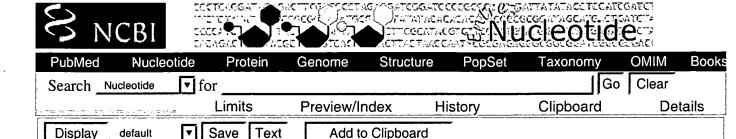
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11



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              A tetrodotoxin-resistant voltage-gated sodium channel expressed by
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              Wood, J.N.
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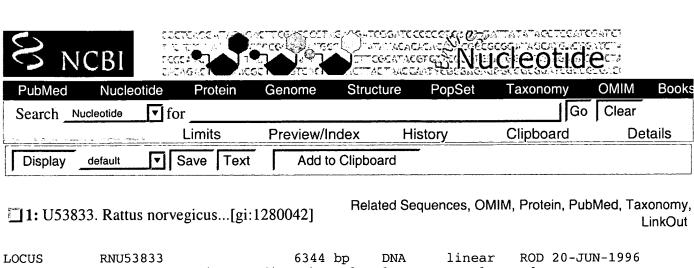
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3 of 4 5/11/02 3:00 PM



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              Stewart, G.R., Sze, P., Hunter, J.C., Eglen, R.M. and Herman, R.C.
    TITLE
              Structure and function of a novel voltage-gated,
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              J. Biol. Chem. 271 (11), 5953-5956 (1996)
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    AUTHORS
              Sangameswaran, L., Delgado, S.G., Fish, L.M. and Herman, R.C.
    TITLE
              Direct Submission
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    JOURNAL
              Neurobiology Unit, Roche Bioscience, 3401, Hillview Avenue, Palo
              Alto, CA 94304, USA
                 (bases 1 to 6344)
  REFERENCE
              Sangameswaran, L.B., Delgado, S.G., Fish, L.M., Koch, B.D.,
    AUTHORS
              Jakeman, L.B., Stewart, G.R., Sze, P., Hunter, J.C., Eglen, R.M. and
              Herman, R.C.
              Additions and corrections to structure and function of a novel
    TITLE
              voltage-gated, tetrodotoxin-resistant sodium channel specific to
              sensory neurons
    JOURNAL
              J. Biol. Chem. 271 (22), 13292-13292 (1996)
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3 of 4 5/11/02 3:00 PM



PubMed	Nucleotide	Protein	Genome	Structure	PopSet	Taxonomy	OMIM	Books
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1: P08104. Sodium channel pr...[gi:116449]

BLink, Related Sequences, PubMed, Taxonomy, LinkOut

LOCUS CIN3_RAT 1951 aa linear ROD 16-OCT-2001 DEFINITION Sodium channel protein, brain III alpha subunit (Voltage-gated sodium channel subtype III).

ACCESSION P08104 PID 9116449

VERSION P08104 GI:116449

DBSOURCE swissprot: locus CIN3_RAT, accession P08104;

class: standard. created: Aug 1, 1988.

sequence updated: Aug 1, 1988. annotation updated: Oct 16, 2001. xrefs: gi: 57210, gi: 57211, gi: 92754

xrefs (non-sequence databases): InterPro IPR002111, InterPro
IPR000636, InterPro IPR001682, InterPro IPR000048, InterPro
IPR001696, Pfam PF00520, Pfam PF00612, PRINTS PR00170, SMART

SM00015

KEYWORDS Ionic channel; Transmembrane; Ion transport; Voltage-gated channel;

Glycoprotein; Repeat; Multigene family.

SOURCE Norway rat.

ORGANISM Rattus norvegicus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

Rattus.

REFERENCE 1 (residues 1 to 1951)

AUTHORS Kayano, T., Noda, M., Flockerzi, V., Takahashi, H. and Numa, S.

TITLE Primary structure of rat brain sodium channel III deduced from the

cDNA sequence

JOURNAL FEBS Lett. 228 (1), 187-194 (1988)

MEDLINE 88137594

REMARK SEQUENCE FROM N.A.

STRAIN=WISTAR

COMMENT

This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. The original entry is available from http://www.expasy.ch/sprot and http://www.ebi.ac.uk/sprot

[FUNCTION] THIS PROTEIN MEDIATES THE VOLTAGE-DEPENDENT SODIUM ION PERMEABILITY OF EXCITABLE MEMBRANES. ASSUMING OPENED OR CLOSED CONFORMATIONS IN RESPONSE TO THE VOLTAGE DIFFERENCE ACROSS THE MEMBRANE, THE PROTEIN FORMS A SODIUM-SELECTIVE CHANNEL THROUGH WHICH NA++ IONS MAY PASS IN ACCORDANCE WITH THEIR ELECTROCHEMICAL GRADIENT.

[SUBUNIT] THE SODIUM CHANNEL CONSISTS OF A LARGE POLYPEPTIDE AND 2-3 SMALLER ONES. THIS SEQUENCE REPRESENTS A LARGE POLYPEPTIDE. [SUBCELLULAR LOCATION] INTEGRAL MEMBRANE PROTEIN.

[DOMAIN] THE SEQUENCE CONTAINS 4 INTERNAL REPEATS, EACH WITH 5 HYDROPHOBIC SEGMENTS (S1,S2,S3,S5,S6) AND ONE POSITIVELY CHARGED SEGMENT (S4). SEGMENTS S4 ARE PROBABLY THE VOLTAGE-SENSORS AND ARE CHARACTERIZED BY A SERIES OF POSITIVELY CHARGED AMINO ACIDS AT

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EVERY THIRD POSITION.
            [SIMILARITY] TO OTHER SODIUM CHANNEL PROTEINS.
            [SIMILARITY] CONTAINS 1 IQ DOMAIN.
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     1921 ydsvtkpdke kfekdkpeke ikgkevrenq k
11
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PubMed	Nucleotide	Protein	Genome	Structure	PopSet	Taxonomy	OMIM	Books
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marama de de la Companya de la Comp	1	Limits	Preview/I	ndex l	History	Clipboard	De	tails
Display	default ▼	Save Text	Add to	o Clipboard	_			A STATE OF THE STA

1: P08104. Sodium channel pr...[gi:116449]

BLink, Related Sequences, PubMed, Taxonomy, LinkOut

LOCUS CIN3_RAT 1951 aa linear ROD 16-OCT-2001

DEFINITION Sodium channel protein, brain III alpha subunit (Voltage-gated

sodium channel subtype III).

ACCESSION P08104 PID g116449

VERSION P08104 GI:116449

DBSOURCE swissprot: locus CIN3_RAT, accession P08104;

class: standard.
created: Aug 1, 1988.

sequence updated: Aug 1, 1988. annotation updated: Oct 16, 2001. xrefs: gi: 57210, gi: 57211, gi: 92754

xrefs (non-sequence databases): InterPro IPR002111, InterPro IPR000636, InterPro IPR001682, InterPro IPR000048, InterPro IPR001696, Pfam PF00520, Pfam PF00612, PRINTS PR00170, SMART

SM00015

KEYWORDS Ionic channel; Transmembrane; Ion transport; Voltage-gated channel;

Glycoprotein; Repeat; Multigene family.

SOURCE Norway rat.

ORGANISM Rattus norvegicus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

Rattus.

REFERENCE 1 (residues 1 to 1951)

AUTHORS Kayano, T., Noda, M., Flockerzi, V., Takahashi, H. and Numa, S.

TITLE Primary structure of rat brain sodium channel III deduced from the

cDNA sequence

JOURNAL FEBS Lett. 228 (1), 187-194 (1988)

MEDLINE 88137594

REMARK SEQUENCE FROM N.A.

STRAIN=WISTAR

COMMENT

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[FUNCTION] THIS PROTEIN MEDIATES THE VOLTAGE-DEPENDENT SODIUM ION PERMEABILITY OF EXCITABLE MEMBRANES. ASSUMING OPENED OR CLOSED CONFORMATIONS IN RESPONSE TO THE VOLTAGE DIFFERENCE ACROSS THE MEMBRANE, THE PROTEIN FORMS A SODIUM-SELECTIVE CHANNEL THROUGH WHICH NA++ IONS MAY PASS IN ACCORDANCE WITH THEIR ELECTROCHEMICAL, GRADIENT.

[SUBUNIT] THE SODIUM CHANNEL CONSISTS OF A LARGE POLYPEPTIDE AND 2-3 SMALLER ONES. THIS SEQUENCE REPRESENTS A LARGE POLYPEPTIDE. [SUBCELLULAR LOCATION] INTEGRAL MEMBRANE PROTEIN.

[DOMAIN] THE SEQUENCE CONTAINS 4 INTERNAL REPEATS, EACH WITH 5
HYDROPHOBIC SEGMENTS (S1,S2,S3,S5,S6) AND ONE POSITIVELY CHARGED
SEGMENT (S4). SEGMENTS S4 ARE PROBABLY THE VOLTAGE-SENSORS AND ARE
CHARACTERIZED BY A SERIES OF POSITIVELY CHARGED AMINO ACIDS AT

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EVERY THIRD POSITION.
            [SIMILARITY] TO OTHER SODIUM CHANNEL PROTEINS.
            [SIMILARITY] CONTAINS 1 IQ DOMAIN.
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3 of 5

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      301 sngtfvnvtm stfnwkdyia ddshfyvldg qkdpllcgng sdagqcpegy icvkagrnpn
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      421 avvamayeeq nqatleeaeq keaefqqmle qlkkqqeeaq avaaasaasr dfsgigglge
      481 llessseask lssksakewr nrrkkrrqre hlegnhradg drfpksesed svkrrsflls
      541 ldgnpltgdk klcsphqsll sirgslfspr rnsktsifsf rgrakdvgse ndfaddehst
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